

	Description
Function	Trainee
Location / Contact	AC Immune SA, EPFL Innovation Park, Building B, 1015 Lausanne <u>careers@acimmune.com</u>
Percentage	100 % - 6 months internship
Reporting Line	Senior Scientist in the Chemistry team
Company Profile	<ul> <li>AC Immune is a clinical stage Swiss biotech company focused on the development of innovative therapeutics and diagnostics for Alzheimer's and other neurodegenerative diseases</li> <li>150+ Employees, 20+ nationalities, IPO in 2016, listed on NASDAQ</li> <li>AC Immune SA is a progressive, equal opportunity employer</li> </ul>
Job description	Trainee position within AC Immune's R&D department, to support the discovery and development of new drugs for the treatment of neurodegenerative and inflammatory diseases.
	AC Immune is seeking a motivated candidate with a strong scientific background in computational biology and biochemistry. The candidate will be contributing to the exploration and implementation of computational methods applied to drug design, with a focus on molecular docking and protein-ligand interaction studies.
	The candidate must keep an excellent record of the <i>in silico</i> experiments and comprehensive documentation using electronic notebooks.
Key Responsibilities	<ul> <li>Execute standard analysis of apo and holo protein structures with small molecules and generate quantitative structure activity relationships and calculate docking scores and potential binding energies.</li> <li>Perform virtual screening and analyze of protein-ligand interactions using docking softwares (e.g. AutoDock Suite and PyRx)</li> <li>Assist in the interpretation and visualization of docking results using computation tools (e.g Pymol and AlphaFold)</li> <li>Analyze, record/document and present the scientific results</li> <li>Collaborate with biologist and chemist and participate in general data management activities with data science team</li> </ul>
Qualifications & Skills	Required:
	<ul> <li>Last year of a master's/engineer degree in biotechnology, computer science, computational biology or similar discipline.</li> <li>Familiarity with structural biological tools such Pymol and Chimera or other molecular visualization programs.</li> <li>Experience with Linux/Unix environments and command-line tools is a plus.</li> <li>Basic knowledge of machine learning or data science frameworks (e.g. scikit-learn) is advantageous but not required.</li> </ul>
	Personal features should include:
	<ul> <li>Team player paired with autonomy and organization skills</li> <li>Good spoken and written English are required</li> </ul>